

Amendments to the Specification:

Please replace the paragraph beginning at page 18, line 9 with the following amended paragraph:

OATP-C genotype data for 3 SNPs Asn130Asp, Pro 155Thr and Val174Ala was utilised to determine the haplotypes. The package SNPHAP (Clayton, David SNPHAPv0.2 2002 <http://www-gene.cimr.cam.ac.uk/clayton/software/snphap.txt> world wide web-gene at [cimr.cam.ac.uk/clayton/software/snphap.txt](http://cimr.cam.ac.uk/clayton/software/snphap.txt)) was used for this analysis. The haplotypes were also predicted using the PHASE package (Stephens, M. 2001 American Journal of Human Genetics 68, 978-989) and were found to give the same predicted haplotypes. The haplotypes were found to be concordant with those reported previously (Tirona 2001). The -118 A>C promoter SNP, at the NF1 binding site, was in strong linkage disequilibrium with the Val174Ala coding variant ( $\Delta=0.3$ ).

Please replace the paragraph beginning on page 18, line 17 with the following rewritten paragraph:

**Table 1 Common haplotypes in the OATP-C gene**

* nomenclature	Amino acid variant(s) on allele	Haplotype frequency (n=79)
*1a	<del>Asp130</del> <u>Asn130</u> , Pro155, Val174	57%
*1b	<del>Asn130</del> <u>Asp130</u> , Pro155, Val174	22%
*5	<del>Asp130</del> <u>Asn130</u> , Pro155, <del>Ala174</del>	2%
*14	<del>Asn130</del> <u>Asp130</u> , Thr155, Val174	7.5%
*15	<del>Asn130</del> <u>Asp130</u> , Pro155, <del>Ala174</del>	11.5%